

GenCore version 5.1.6
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OM protein --protein search, using sw model

Run on: November 12, 2003, 15:41:34 ; Search time 9.84282 Seconds
(without alignments)
640.499 Million cell updates/sec

Title: US-09-963-347B-2

Perfect score: 769

Sequence: 1 MGCPRMFPALLYVLSVSR.....KRRKRKVTNNKLEQVSQL 149

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 328717 seqs, 42310858 residues

Total number of hits satisfying chosen parameters: 328717

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Issued Patents AA:*

- 1: /cgn2_6/prodata/2/iaa/5A COMB.pep.*
- 2: /cgn2_6/prodata/2/iaa/5B COMB.pep.*
- 3: /cgn2_6/prodata/2/iaa/6A COMB.pep.*
- 4: /cgn2_6/prodata/2/iaa/6B COMB.pep.*
- 5: /cgn2_6/prodata/2/iaa/PCITUS COMB.pep.*
- 6: /cgn2_6/prodata/2/iaa/backfile1.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | % Match | Length | ID | Description |
|------------|-------|---------|--------|----|--------------------|
| 1 | 707 | 91.3 | 159 | 4 | US-09-852-391-2 |
| 2 | 81.5 | 10.6 | 674 | 4 | US-09-328-352-6455 |
| 3 | 73 | 9.5 | 1461 | 2 | US-08-993-228-10 |
| 4 | 72.5 | 9.4 | 1465 | 4 | US-09-627-986-1 |
| 5 | 71.5 | 9.3 | 1394 | 3 | US-09-213-053-2 |
| 6 | 71.5 | 9.3 | 2184 | 4 | US-09-417-485D-6 |
| 7 | 71 | 9.2 | 154 | 1 | US-08-446-908-2 |
| 8 | 71 | 9.2 | 154 | 1 | US-08-231-205A-2 |
| 9 | 71 | 9.2 | 154 | 2 | US-08-871-161-2 |
| 10 | 71 | 9.2 | 667 | 4 | US-09-328-352-5747 |
| 11 | 70 | 9.1 | 1155 | 1 | US-08-286-889-46 |
| 12 | 70 | 9.1 | 1155 | 1 | US-08-485-618-46 |
| 13 | 70 | 9.1 | 1155 | 1 | US-08-362-652-46 |
| 14 | 70 | 9.1 | 1155 | 2 | US-08-605-672-46 |
| 15 | 70 | 9.1 | 1155 | 2 | US-08-482-293A-46 |
| 16 | 70 | 9.1 | 1155 | 2 | US-08-943-363-46 |
| 17 | 70 | 9.1 | 1155 | 3 | US-09-193-043-46 |
| 18 | 70 | 9.1 | 1161 | 1 | US-08-688-307A-46 |
| 19 | 70 | 9.1 | 1161 | 1 | US-08-485-618-53 |
| 20 | 70 | 9.1 | 1161 | 2 | US-08-362-652-53 |
| 21 | 70 | 9.1 | 1161 | 2 | US-08-605-672-53 |
| 22 | 70 | 9.1 | 1161 | 2 | US-08-482-293A-53 |
| 23 | 70 | 9.1 | 1161 | 2 | US-08-943-363-53 |
| 24 | 70 | 9.1 | 1161 | 3 | US-09-193-043-53 |
| 25 | 70 | 9.1 | 1161 | 3 | US-09-688-307A-53 |
| 26 | 69 | 9.0 | 1589 | 3 | US-09-356-952-4 |
| 27 | 68 | 8.8 | 440 | 3 | US-08-430-286A-9 |

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| 28 | 68 | 8.8 | 2285 | 4 | US-09-308-375-2 | Sequence 2, Appli |
| 29 | 67.5 | 8.8 | 354 | 4 | US-09-198-452A-317 | Sequence 217, App |
| 30 | 67 | 8.7 | 154 | 4 | US-09-252-991A-27368 | Sequence 27368, A |
| 31 | 67 | 8.7 | 411 | 1 | US-07-937-609-21 | Sequence 21, Appl |
| 32 | 67 | 8.7 | 411 | 3 | US-08-029-170-21 | Sequence 21, Appl |
| 33 | 67 | 8.7 | 459 | 1 | US-08-220-151-12 | Sequence 12, Appl |
| 34 | 67 | 8.7 | 459 | 1 | US-08-220-151-14 | Sequence 12, Appl |
| 35 | 67 | 8.7 | 459 | 1 | US-08-413-118-12 | Sequence 12, Appl |
| 36 | 67 | 8.7 | 459 | 1 | US-08-413-118-14 | Sequence 14, Appl |
| 37 | 67 | 8.7 | 459 | 3 | US-08-473-446-12 | Sequence 12, Appl |
| 38 | 67 | 8.7 | 459 | 3 | US-08-473-446-14 | Sequence 14, Appl |
| 39 | 67 | 8.7 | 459 | 3 | US-09-213-053-6 | Sequence 6, Appli |
| 40 | 66.5 | 8.6 | 531 | 5 | PCT-US92-00282-6 | Sequence 6, Appli |
| 41 | 66 | 8.6 | 451 | 3 | US-08-430-286A-10 | Sequence 10, Appl |
| 42 | 66 | 8.6 | 782 | 1 | US-07-725-083-2 | Sequence 2, Appli |
| 43 | 66 | 8.6 | 782 | 3 | US-08-669-286-10 | Sequence 10, Appl |
| 44 | 66 | 8.6 | 782 | 3 | US-09-469-253-10 | Sequence 10, Appl |
| 45 | 66 | 8.6 | 782 | 3 | US-09-642-146-10 | Sequence 10, Appl |

ALIGNMENTS

RESULT 1
US-09-852-391-2
; Sequence 2, Application US/098523391
; Patent No. 6555520
; GENERAL INFORMATION:
; APPLICANT: Sims, John
; APPLICANT: Lyman, Stewart
; APPLICANT: Armstrong, Allison
; APPLICANT: McKenna, Hilary
; TITLE OF INVENTION: Human TSLP DNA and Polypeptides
; FILE REFERENCE: 03260.0087-00304/2881-WO
; CURRENT APPLICATION NUMBER: US/09/852,391
; CURRENT FILING DATE: 2001-05-09
; PRIOR APPLICATION NUMBER: 60/108,452
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: Patent In Ver. 2.0
; SEQ ID NO 2
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-852-391-2

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| Best Local Similarity | 96.5% | Fred. No. 2.5e-75; | | |
| Matches 139; | Conservative 0; | Mismatches 5; | Indels 0; | Gaps 0; |
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| Db | 1 | MPPFALLVLSVSPKFIQLQVGLVLYDFNCDFFKIKAAVLTISKDLITYMSGTKS | 60 | |
| QY | 66 | TEFNNTVSCSNRPCHLTETIQSLTFNPNRVRSLAKEMFAMTKAALAIWCPGYSTQINA | 125 | |
| Db | 61 | TEFNNTVSCSNRPCHLTETIQSLTFNPTAGCAGSLAKEMFAMTKAALAIWCPGYSTQINA | 120 | |
| QY | 126 | TOAMKRRKRKVTNNKLEQVSQL | 149 | |
| Db | 121 | TOAMKRRKRKVTNNKLEQVSQL | 144 | |

RESULT 2
US-09-328-352-6455
; Sequence 6455, Application US/09328352
; Patent No. 6562958
; GENERAL INFORMATION:
; APPLICANT: Gary L. Berton et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
; FILE REFERENCE: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
; CURRENT APPLICATION NUMBER: GTC99-03PA
; CURRENT APPLICATION NUMBER: US/09/328,352

| Result No. | Score | Query | | DB | ID | Description |
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| 3 | 33.4 | 7.0 | 1664976 | 4 | US-08-916-421B-1 | Sequence 1, Appli |
| 4 | 31.8 | 6.6 | 1881 | 5 | PTC-US94-09752-2 | Sequence 2, Appli |
| 5 | 31 | 6.5 | 1734 | 4 | US-09-288-143-42 | Sequence 4, Appli |
| 6 | 31 | 6.5 | 2460 | 4 | US-09-328-532-524 | Sequence 524, App |
| 7 | 31 | 6.5 | 1830121 | 4 | US-09-557-884-1 | Sequence 1, Appli |
| 8 | 31 | 6.5 | 1830121 | 4 | US-09-643-990A-1 | Sequence 1, Appli |
| 9 | 30.4 | 6.3 | 1218 | 4 | US-09-252-991A-14566 | Sequence 14566, A |
| 10 | 30.4 | 6.3 | 4403765 | 3 | US-09-103-840A-2 | Sequence 2, Appli |
| 11 | 30.4 | 6.3 | 4411529 | 3 | US-09-103-840A-1 | Sequence 1, Appli |
| 12 | 30.2 | 6.3 | 690 | 4 | US-09-404-879A-321 | Sequence 321, App |
| 13 | 30 | 6.2 | 231 | 1 | US-08-026-145-5 | Sequence 5, Appli |
| 14 | 30 | 6.2 | 234 | 1 | US-08-446-646-10 | Sequence 10, Appl |
| 15 | 30 | 6.2 | 450 | 1 | US-08-446-646-4 | Sequence 4, Appli |
| 16 | 29.6 | 6.2 | 2373 | 3 | US-08-817-707-5 | Sequence 5, Appli |
| 17 | 29.6 | 6.2 | 38844 | 4 | US-09-734-675-3 | Sequence 3, Appli |
| 18 | 29.6 | 6.2 | 1664976 | 4 | US-08-916-421B-1 | Sequence 1, Appli |
| 19 | 29.6 | 6.2 | 1830121 | 4 | US-09-557-884-1 | Sequence 1, Appli |
| 20 | 29.6 | 6.2 | 1830121 | 4 | US-09-643-990A-1 | Sequence 1, Appli |
| 21 | 29.4 | 6.1 | 998 | 2 | US-07-885-089B-5 | Sequence 5, Appli |
| 22 | 29.4 | 6.1 | 1282 | 4 | US-08-858-207A-153 | Sequence 153, App |
| 23 | 29.2 | 6.1 | 1001 | 4 | US-09-671-317-412 | Sequence 412, App |
| 24 | 29.2 | 6.1 | 1073 | 4 | US-09-481-049-6 | Sequence 6, Appli |
| 25 | 29.2 | 6.1 | 1976 | 4 | US-09-356-806-112 | Sequence 112, App |
| 26 | 29.2 | 6.1 | 2312 | 4 | US-09-356-806-114 | Sequence 114, App |
| 27 | 29.2 | 6.1 | 3212 | 3 | US-08-697-954-1 | Sequence 1, Appli |

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OM protein - protein search, using sw model

Run on: November 12, 2003, 15:41:34 ; Search time 10.5034 Seconds
(without alignments)
640.499 Million cell updates/sec

Title: US-09-963-347B-4
Perfect score: 823
Sequence: 1 MPPFALLVLSVSPKIFIL.....QVSQLGLWRNRPRLKQQ 159

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 328717 seqs, 4231058 residues

Total number of hits satisfying chosen parameters: 328717

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued Patents AA:*
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2: /cgm2_6/ptodata/2/iaa/5B COMB.pcp.*
3: /cgm2_6/ptodata/2/iaa/6A COMB.pcp.*
4: /cgm2_6/ptodata/2/iaa/6B COMB.pcp.*
5: /cgm2_6/ptodata/2/iaa/PCTUS_COMB.pcp.*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match | Length | DB ID | Description |
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| 1 | 823 | 100.0 | 159 | 4 | US-09-852-391-2 |
| 2 | 74.5 | 9.1 | 674 | 4 | US-09-328-352-6455 |
| 3 | 73 | 8.9 | 1155 | 1 | US-08-286-889-46 |
| 4 | 73 | 8.9 | 1155 | 1 | US-08-485-618-46 |
| 5 | 73 | 8.9 | 1155 | 1 | US-08-362-652-46 |
| 6 | 73 | 8.9 | 1155 | 2 | US-08-605-672-46 |
| 7 | 73 | 8.9 | 1155 | 2 | US-08-482-293A-46 |
| 8 | 73 | 8.9 | 1155 | 2 | US-08-943-363-46 |
| 9 | 73 | 8.9 | 1155 | 3 | US-09-193-043-46 |
| 10 | 73 | 8.9 | 1155 | 4 | US-09-688-307A-46 |
| 11 | 73 | 8.9 | 1161 | 1 | US-08-485-618-53 |
| 12 | 73 | 8.9 | 1161 | 1 | US-08-362-652-53 |
| 13 | 73 | 8.9 | 1161 | 2 | US-08-605-672-53 |
| 14 | 73 | 8.9 | 1161 | 2 | US-08-482-293A-53 |
| 15 | 73 | 8.9 | 1161 | 2 | US-08-943-363-53 |
| 16 | 73 | 8.9 | 1161 | 3 | US-09-193-043-53 |
| 17 | 73 | 8.9 | 1161 | 4 | US-09-688-307A-53 |
| 18 | 73 | 8.9 | 1461 | 2 | US-08-993-228-10 |
| 19 | 71 | 8.6 | 404 | 4 | US-09-198-452A-415 |
| 20 | 69.5 | 8.4 | 531 | 5 | PCT-US92-00282-6 |
| 21 | 69.5 | 8.4 | 711 | 3 | US-08-949-588-2 |
| 22 | 69.5 | 8.4 | 1394 | 3 | US-09-213-053-2 |
| 23 | 68.5 | 8.3 | 465 | 4 | US-08-627-986-1 |
| 24 | 68 | 8.3 | 328 | 4 | US-09-180-827-7 |
| 25 | 68 | 8.3 | 1589 | 3 | US-09-356-952-4 |
| 26 | 68 | 8.3 | 2285 | 4 | US-09-308-375-2 |
| 27 | 67 | 8.1 | 154 | 4 | US-09-252-991A-27368 |

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| 28 | 66 | 8.0 | 314 | 4 | US-09-328-352-4934 |
| 29 | 66 | 8.0 | 321 | 3 | US-08-673-814-2 |
| 30 | 66 | 8.0 | 321 | 4 | US-09-115-824-2 |
| 31 | 66 | 8.0 | 366 | 3 | US-08-975-762-53 |
| 32 | 66 | 8.0 | 366 | 3 | US-09-295-028-53 |
| 33 | 66 | 8.0 | 366 | 4 | US-09-106-582-53 |
| 34 | 66 | 8.0 | 667 | 4 | US-09-328-352-5747 |
| 35 | 66 | 8.0 | 1293 | 4 | US-09-170-496B-292 |
| 36 | 66 | 8.0 | 2183 | 3 | US-08-746-111-5 |
| 37 | 65.5 | 8.0 | 127 | 3 | US-08-946-026-4 |
| 38 | 65.5 | 8.0 | 321 | 4 | US-08-948-276-4 |
| 39 | 65.5 | 8.0 | 612 | 1 | US-08-344-695-2 |
| 40 | 65 | 7.9 | 154 | 1 | US-08-446-308-2 |
| 41 | 65 | 7.9 | 154 | 1 | US-08-231-205A-2 |
| 42 | 65 | 7.9 | 154 | 2 | US-08-871-161-2 |
| 43 | 65 | 7.9 | 427 | 1 | US-08-361-920-25 |
| 44 | 65 | 7.9 | 427 | 1 | US-08-479-939-25 |
| 45 | 65 | 7.9 | 427 | 1 | US-08-483-432-25 |

ALIGNMENTS

RESULT 1
US-09-852-391-2
; Sequence 2, Application US/09852391
; Patent No. 655520
; GENERAL INFORMATION:

; APPLICANT: Sims, John
; APPLICANT: Lyman, Stewart
; APPLICANT: Armstrong, Allison
; APPLICANT: McKenna, Hilary
; TITLE OF INVENTION: Human TSLP DNA and Polypeptides
; FILE REFERENCE: 03260.0087-00304/2881-WO
; CURRENT APPLICATION NUMBER: US/09/852,391
; PRIOR FILING DATE: 2001-05-09
; PRIOR APPLICATION NUMBER: 60/108,452
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: Patent In Ver. 2.0
; SEQ ID NO 2
; LENGTH: 159
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-852-391-2

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| Matches 159; | Conservative 0; | | | Gaps 0; |
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| QY | 61 | TEFNNTVSCSRPCHLTIQSLTENPTAGCASAKENFAMKTKAALAIWCPGVSETQINA | 120 | |
| Db | 61 | TEFNNTVSCSRPCHLTIQSLTENPTAGCASAKENFAMKTKAALAIWCPGVSETQINA | 120 | |
| QY | 121 | TQAMKRRKRVTKNKCLEQVSQLQGLWRNRPRLKQQ | 159 | |
| Db | 121 | TQAMKRRKRVTKNKCLEQVSQLQGLWRNRPRLKQQ | 159 | |

RESULT 2
US-09-328-352-6455
; Sequence 6455, Application US/09328352
; Patent No. 6562958
; GENERAL INFORMATION:

; APPLICANT: Gary L. Breton et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
; TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: GTC99-03PA
; CURRENT APPLICATION NUMBER: US/09/328,352

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OM protein - protein search, using sw model

Run on: November 12, 2003, 15:41:34 ; Search time 8.65376 Seconds
(without alignments)
640.499 Million cell updates/sec

Title: US-09-963-347B-4_COPY_1_131

Perfect score: 675
Sequence: 1 MFPFALLVLSVSRKIFIL.....GYSETQINATQAMKRRKK 131

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 328717 seqs, 42310858 residues

Total number of hits satisfying chosen parameters: 328717

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

- Issued Patents AA:*
- 1: /cgn2_6/ptodata/2/iaa/5A COMB.pep.*
 - 2: /cgn2_6/ptodata/2/iaa/5B COMB.pep.*
 - 3: /cgn2_6/ptodata/2/iaa/6A COMB.pep.*
 - 4: /cgn2_6/ptodata/2/iaa/6B COMB.pep.*
 - 5: /cgn2_6/ptodata/2/iaa/PCTUS COMB.pep.*
 - 6: /cgn2_6/ptodata/2/iaa/backfiles1.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match | Length | DB ID | Description |
|------------|-------|-------------|--------|-------|----------------------|
| 1 | 675 | 100.0 | 159 | 4 | US-09-852-391-2 |
| 2 | 73 | 10.8 | 1155 | 1 | US-08-286-889-46 |
| 3 | 73 | 10.8 | 1155 | 1 | US-08-485-618-46 |
| 4 | 73 | 10.8 | 1155 | 1 | US-08-362-652-46 |
| 5 | 73 | 10.8 | 1155 | 2 | US-08-605-672-46 |
| 6 | 73 | 10.8 | 1155 | 2 | US-08-482-293A-46 |
| 7 | 73 | 10.8 | 1155 | 2 | US-08-943-363-46 |
| 8 | 73 | 10.8 | 1155 | 3 | US-09-193-043-46 |
| 9 | 73 | 10.8 | 1155 | 4 | US-09-688-307A-46 |
| 10 | 73 | 10.8 | 1161 | 1 | US-08-485-618-53 |
| 11 | 73 | 10.8 | 1161 | 1 | US-08-362-652-53 |
| 12 | 73 | 10.8 | 1161 | 2 | US-08-605-672-53 |
| 13 | 73 | 10.8 | 1161 | 2 | US-08-482-293A-53 |
| 14 | 73 | 10.8 | 1161 | 2 | US-08-943-363-53 |
| 15 | 73 | 10.8 | 1161 | 3 | US-09-193-043-53 |
| 16 | 73 | 10.8 | 1161 | 4 | US-09-688-307A-53 |
| 17 | 73 | 10.8 | 1461 | 2 | US-08-993-228-10 |
| 18 | 69.5 | 10.3 | 531 | 5 | PCT-US92-00282-6 |
| 19 | 68 | 10.1 | 1589 | 3 | US-09-356-952-4 |
| 20 | 67 | 9.9 | 2285 | 4 | US-09-308-375-2 |
| 21 | 66 | 9.8 | 366 | 3 | US-08-975-762-53 |
| 22 | 66 | 9.8 | 366 | 3 | US-09-295-028-53 |
| 23 | 66 | 9.8 | 366 | 4 | US-09-106-582-53 |
| 24 | 65.5 | 9.7 | 321 | 4 | US-08-948-276-4 |
| 25 | 65.5 | 9.7 | 465 | 4 | US-09-627-986-1 |
| 26 | 65 | 9.6 | 154 | 4 | US-09-252-991A-27368 |
| 27 | 65 | 9.6 | 589 | 1 | US-07-668-648-6 |

Sequence 6, Appli
Sequence 6, Appli
Sequence 2, Appli
Sequence 6, Appli
Sequence 4, Appli
Sequence 4455, Ap
Sequence 29, Appli
Sequence 15, Appli
Sequence 17, Appli
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Sequence 3795, Ap
Sequence 121, App
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Sequence 505, App

ALIGNMENTS

RESULT 1
US-09-852-391-2
; Sequence 2, Application US/09852391
; Patent No. 655520
; GENERAL INFORMATION:

; APPLICANT: Sims, John
; APPLICANT: Lyman, Stewart
; APPLICANT: Armstrong, Allison
; APPLICANT: McKenna, Hilary
; TITLE OF INVENTION: Human TSLP DNA and Polypeptides
; FILE REFERENCE: 03260-0087-00304/2881-WO
; CURRENT APPLICATION NUMBER: US/09/852.391
; PRIOR FILING DATE: 2001-05-09
; PRIOR APPLICATION NUMBER: 60/108,452
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: Patent In Ver. 2.0
; SEQ ID NO 2

; LENGTH: 159
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-852-391-2

Query Match 100.0%; Score 675; DB 4; Length 159;
Best Local Similarity 100.0%; Pred. No. 2.3e-73;
Matches 131; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MFPFALLVLSVSRKIFILQVLGLVLTPTNCDPEKIKAAVLTSTISKDLITYMSGTKS 60

Db 1 MFPFALLVLSVSRKIFILQVLGLVLTPTNCDPEKIKAAVLTSTISKDLITYMSGTKS 60

QY 61 TEFNNTVSCSRNPCHLTETIQLTNPAGCASLAKENPAMTKAALAIWCPGSETQINA 120

Db 61 TEFNNTVSCSRNPCHLTETIQLTNPAGCASLAKENPAMTKAALAIWCPGSETQINA 120

QY 121 TQAMKRRKK 131

Db 121 TQAMKRRKK 131

RESULT 2

US-08-286-889-46

; Sequence 46, Application US/08286889

; Patent No. 5470953

; GENERAL INFORMATION:

; APPLICANT: Gallatin, W. Mich

; APPLICANT: Van der Vieren, Monica

; TITLE OF INVENTION: NO. 5470953e1 Human 2 Integrin Alpha Subunit

; NUMBER OF SEQUENCES: 51

; CORRESPONDENCE ADDRESS: